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## **AMENDMENT TO THE CLAIMS:**

This listing of the claims will replace all prior versions, and listings, of the claims in the application.

Claims 1-38. (canceled)

Claim 39. (new) A chimeric gene capable of causing an increased level of lysine in seeds obtained from a transformed plant, the chimeric gene comprising:

- a) an isolated nucleic acid fragment comprising a nucleic acid sequence which is useful in antisense inhibition or sense suppression of endogenous lysine ketoglutarate reductase/saccharopine dehydrogenase activity in a plant or plant cell wherein said isolated nucleic acid fragment comprises all or a part of the nucleic acid sequence encoding a plant lysine ketoglutarate reductase/saccharopine dehydrogenase, said part being sufficient for use in antisense inhibition or sense suppression; and
- b) at least one regulatory sequence operably linked to said fragment.
  Claim 40. (new) A plant comprising the chimeric gene of claim 39 in its genome.
  - Claim 41. (new) Seed obtained from the plant of claim 40.
- Claim 42. (new) A method for increasing lysine content in a plant seed which comprises:
  - (a) transforming plant cells with the chimeric gene of claim 39;
- (b) regenerating fertile mature plants from the transformed plant cells obtained from step (a) under conditions suitable to obtain seeds;
- (c) screening progeny seed of step (b) for increased lysine content; and
- (d) selecting those lines whose seeds have increased lysine content.
  - Claim 43. (new) Seed obtained by the method of claim 42.
- Claim 44. (new) A chimeric gene capable of causing an increased level of lysine in seeds obtained from a transformed corn plant, the chimeric gene comprising:

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- a) an isolated nucleic acid fragment comprising a nucleic acid sequence which is useful in antisense inhibition or sense suppression of endogenous lysine ketoglutarate reductase/saccharopine dehydrogenase activity in a corn plant or corn plant cell wherein said isolated nucleic acid fragment comprises all or a part of the nucleic acid sequence encoding a corn plant lysine ketoglutarate reductase/saccharopine dehydrogenase, said part being sufficient for use in antisense inhibition or sense suppression; and
  - b) at least one regulatory sequence operably linked to said fragment.
    Claim 45. (new) A corn plant comprising the chimeric gene of claim 44 in its genome.
    - Claim 46. (new) Seed obtained from the corn plant of claim 45.
- Claim 47. (new) A method for increasing lysine content in a corn plant seed which comprises:
- (a) transforming corn plant cells with the chimeric gene of claim 44;
- (b) regenerating fertile mature plants from the transformed corn plant cells obtained from step (a) under conditions suitable to obtain seeds;
- (c) screening progeny seed of step (b) for increased lysine content; and
- (d) selecting those lines whose seeds have increased lysine content.
  - Claim 48. (new) Seed obtained by the method of claim 47.
- Claim 49. (new) A chimeric gene capable of causing an increased level of lysine in seeds obtained from a transformed corn plant, the chimeric gene comprising:
- a) an isolated nucleic acid fragment comprising a nucleic acid sequence which is useful in antisense inhibition or sense suppression of endogenous lysine ketoglutarate reductase/saccharopine dehydrogenase activity in a corn plant or plant cell wherein said isolated nucleic acid fragment comprises all or a part of the nucleic acid sequence of SEQ ID NO:120, said part being sufficient for use in antisense inhibition or sense suppression; and
  - b) at least one regulatory sequence operably linked to said fragment.

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Claim 50. (new) A plant comprising the chimeric gene of claim 49 in its genome.

Claim 51. (new) Seed obtained from the plant of claim 50.

Claim 52. (new) A method for increasing lysine content in a plant seed which comprises:

- (a) transforming plant cells with the chimeric gene of claim 49;
- (b) regenerating fertile mature plants from the transformed corn plant cells obtained from step (a) under conditions suitable to obtain seeds;
- (c) screening progeny seed of step (b) for increased lysine content; and
- (d) selecting those lines whose seeds have increased lysine content.

Claim 53. (new) Seed obtained by the method of claim 52.